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A26411 21:904967
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Wallach,D. and Brakebusch,C.
Multimers of the soluble forms of TNF receptors, and pharmaceutical compositions containing them Patent: EP 0526905-A 1 10-FEB-1993;
YEDA RESEARCH AND DEVELOPMENT CO. LTD
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YEDA RESEARCH AND DEVELOPMENT COMPANY LIMITED
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of psv-TBP.
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AX404882/c
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(campbell,R.K., Jameson,B.A) a
(campbell heterodimeric protein
(patent: US 6193972-A 19 27-FE
                                                                                                                                                                                                                                                                                                                                                                                             Sequence
AR134771
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ARI 31319
ARI 31319.1 GI:14120222
                                                                                                                                                                                                                                                                                              1 (bases 1 to 21)
Campbell,R.K., Jameson,B.A.
DNA encoding a hybrid hetero
Patent: US 6194177-A 19 27-F
Pfizenmaier, K., Wuest, T., Fusion protein from antib
                                        synthetic construct.
synthetic construct
                                                                                                                                                                                                                                                                                                                                                  Unknown
                                                                     AX404882.1
                                                                                  Sequence 15 from Patent AX404882
                                                                                                                                                                                                                                                                                                                                                             Unknown.
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                             artificial sequences.
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Pred. No. 5.6
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Lbody cytokine-c
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s 6193972.
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5.6e+05;
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5.6e+05;
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                                                                                       Unclassified.

1 (bases 1 to
Masucci, M.G.
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AR052978
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A57512
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Location/Qualifiers
                                                                                                                         Unknown
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                                                                                                                                                                                                                                                                                                                                                                                                           Patent: WO 9632483-A 4 17-OCT-1996;
MASUCCI MARIA GRAZIA (SE)
Other publication AU 5284296 961030.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             unidentified
                                                                             Glycine-containing sequences conferring
                                                                                                                                      Unknown
                                                                                                                                                           AR052978.1
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Masucc1, M.G.
IMMUNE-EVADING PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unclassified
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/note="Primer 6 fuer d
/norte="Primer 6 fuer d
TNFR1-Fragments"
9 c 10 g
                                        Location/Qualifiers
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/db_xref-"taxon:32644"
14 c 2 g
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from Patent W09632483
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Pred. No.
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                                                      10-NOV-1998;
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                                        A57514 24 bp. sequence 6 from Patent W09632483. A57514
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synthetic construct
artificial sequences
Location/Qualifiers
unidentified. unidentified
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Similarity 91.7%;
22. Conservative
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ent: US 5633145-A 15 27-MAY-1997;
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8 c 10 g
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/db_xref="taxon:32630"
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Pred. No. 6.
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MASUCCI,M.G.

IMMONE-EVADING PROTEINS.

Patent: WO 9632483-A 10 17-OCT-1996;

MASUCCI MARKIA GRAKIA (SE)

Other publication AU 5284296 961030.

Location/Qualifiers
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Sequence 10
AR052980
                                                                                                                                                  24 bp
Sequence 10 from Patent WO9632483.
A57518
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MASUCCI MARIA GRAZIA (SE)
Other publication AU 5284296 961030.
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                                                        AAA95191
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Primer for TPO/hCG
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Multimerisation of
Multimerisation of
PCR primer used to
DNA encoding a HRG
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Human 55kDa tumour
Human 55kD TNFBP
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RESULT 1 AAA95191/c ID AAA95191 XX AC AAA95191 XX DT 12-JAN-2 XX FEVETSE XX TNFR1; t KW TNFR1; t KW TUMOUT; XX CS HOMO SAP XX OS HOMO SAP XX	444 W410	4448878655555555555555555555555555555555
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12-SEP-1989;
08-MAR-1990;
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                                                                                                                   Brockhaus M,
Schlaeger E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rolynucleotides comprising polymorphic variants of a referent for tumous necrosis factor receptor 1 (TNFR1), useful for stubiological function of TNFR1 and identifying drugs targeting protein for treating disorders -
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anti-inflammatory;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          apoptosis related disorders and bacterial infection.
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90CH-0000746.
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55 kD

TNFBP extracellular fragment PCR

primer

N

This invention

(in)soluble

describes novel homogeneous insoluble proteins (I), ole fragments (Ia) and their salts that can bind tumour

Example 11;

Page 16;

25pp;

German

invention have antiinflammatory, immunosuppressive, antibacterial, antiprotozoal activity. (I), and related recombinant proteins, are used to treat diseases mediated by TNF, e.g. shock in cases of meningococal sepsis; development of autoimmune glomerulonephritis and cerebral malaria. Also (I), or antibodies specific for them, are used for diagnostic determination of TNF in body fluids, for affinity purification of TNF and for identifying (ant)agonists of TNF. This sequence represents a PCR primer used in the amplification of the human 55 kD TNFBP described

This invention describes novel insoluble proteins (I), also their (in)soluble fragments and pharmaceutically acceptable salts, able tumor necrosis factor (TNF) and in homogeneous form. The products

e t

bind the

Example (TNF),

Page 16;

German.

homogeneous,

is, insoluble proteins that bind tumor necrosis factor for treating TNF-mediated disorders, e.g. inflammation

2001-559312/63

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           glomerulonephritis, cerebral malaria, immune responses and inflammation), (11) to purify TNF, (111) to identify TNF (ant)agonists and (1v) for diagnostic determination of TNF in body fluids. Antibodies raised against (I) are used for affinity purification of (I). This sequence represents a PCR primer used in the amplification of the TNF binding protein of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               necrosis factor (TNF). The products of the invention have anti-inflammatory and antimalarial activity. (I) and (Ia) are to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock, to treat diseases in which TNF is involved (e.g. septic shock).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EP1132471-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    autoimmune glomerulonephritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               insoluble protein; antiinfl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tumor necrosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HOFFMANN LA ROCHE & CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001EP-0108117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dembic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    89CH-000319.
90CH-0000746.
90CH-0001347.
90EP-0116707.
99EP-0100703.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            factor binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gentz
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lesslauer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein; TNFBP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ę
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cerebral malaria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            treatment;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antibacterial;
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             δÃ
                              Query Match
Best Local S
Matches 21
                                                             Sequence
                                                                                        protein encoded by the fusion gene amplified by the present sequence, significantly increases the biological activity of the present sequence.
                                                                            sequence, significantly increases the biological beautiful to the requirement the number of injections needed.
               142 ACTGAGGACTCAGGCACCACA 162
21
                              Similarity 100
21; Conservative
                                                              21
                                                             BP;
                                                              N
                                                              ?
                                                              5 C; 7
                                     100.0%;
                              Score 21; DB; Pred. No. 1.5
                               0
                                                              G; 7
                                                              T; 0
                                                              other;
                              DB 18; 1 . 1.5e+04; ches 0;
                                              Length 21;
                               Indels
                                                                                            of the
                               0
                               0;
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RESULT 5

Query Match Best Local S Matches 24

Similarity

3.6%;

Score 21; Pred. No.

DB 24; 1.7e+04; other;

Length 29;

Conservative

0

Mismatches

Indels

0 Gaps

0

Sequence 29

BP; 3

ð,

9 C; 10 G; 7 T; 0 the disclosure

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Matches 251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT94UI/ Standard; DNA;
                                          A novel fusion protein comprises 2 dimer forming co-expressed amino acid sequences, each consisting of a homodimeric or heterodimeric receptor chain or ligand, with ligand-receptor binding activity, bound directly or via a peptide linker to a subunit of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 29
heterodimeric protein hormone capable of forming a heterodimer with
the hormone's other subunits. The fusion protein, e.g. the
thrombopoletin (TPO)/human chorionic gonadotrophin (hCG) fusion
                                                                                                                                          Hybrid dimeric protein comprising two co-expressed units - expands on receptor or ligand and a subunit of a heterodimeric hormone, especially FSH, for inducing follicular maturation
                                                                                                                                                                                                                                                                                                                                                                                                                                             CG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Primer for TPO/hCG fusion gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT94017;
                                                                                                                 Example;
                                                                                                                                                                                                                                                        (ISTF ) ARS APPLIED RES SYSTEMS HOLDING
                                                                                                                                                                                                                                                                                      20-FEB-1996;
                                                                                                                                                                                                                                                                                                                20-FEB-1997;
                                                                                                                                                                                                                                                                                                                                            21-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                       WO9730161-A1
                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             143 CTGAGGACTCAGGCACCACAGTGCTGT 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29 CTGAGGACTCAGGCACCACAGAGCTCT 3
                                                                                                                                                                                                                                                                                                                                                                                                                                           on protein; thrombopoietin; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  S本1larity
                                                                                                              Page 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                      96US-0011936.
                                                                                                                                                                                                                                                                                                                97WO-US02315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 A; 7 C; 9 G; 8 T; 0 other;
                                                                                                              60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.18;
                                                                                                                                                                                                                              SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
                                                                                                                                                                                                                              Jameson BA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 23.8;
Pred. No. 2.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                           TPO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.6e+03;
                                                                                                                                                                                                                                                          3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         chorionic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         gonadotrophin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
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ABA99921/c
ID ABA99921 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cytostatic; immunomodulator
gene therapy; scFv antibody
solid tumour; angiogenesis;
                                                      only in target tissue, resulting in a high local concentration, and activity is potentiated by co-activation of receptors. This sequence represents a PCR primer for the amplification of the human TNFR1 fracused in the construction of the TNF-selectokine W24 and W33
                                                                                                                                  and/or pathological angiogenesis, also generally for treating infections and metabolic diseases. (I) are prodrug forms of R2 that have unacceptable toxicity when administered systemically (specifically tumour necrosis factor) and allow these compounds to be administered safely with retention of, or even increase in, therapeutic activity. R2 is released only in target tissue, resulting in a high local concentration, and
                                                                                                                                                                                                                                                                          mutant form of the tumour necrosis factor (TNF) precursor protein, a region with a proteolytic cleavage site, and human TNF receptor-1 fragment, and with trypsin (activator) for 5 minutes. After 16 hours, cell viability was determined by MTT staining. Activated W24 had LD50 about 0.5 ng/ml, comparable with that for wild-type TNF and 4000 times higher than for uncleaved W24. (I), also nucleic acids encoding them and related vectors, are useful particularly for treating solid tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polypeptide prodrug, useful e.g. for treating tumors, contains targeting region, active agent and attached inhibitor that is proteolytically cleaved in target cells -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pfizenmaier K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-SEP-2000; 2000DE-1045592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-SEP-2001; 2001WO-EP10730
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               инд00222833-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prodrug; TNF; tumour necrosis factor; selectokine; chimeric; W24; W33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human INFR1 PCR primer SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABA99921;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specific target molecule, a region (R3) that has a processing site and a region (R4) that inhibits the activity of R2, by intramolecular bonding and/or interaction. The products of the invention have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             macromolecule on a cell surface and/or a component of the extracellul matrix, peptide linker, a region (R2) with biological activity for a specific target molecule, a region (R3) that has a processing site an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes a novel polypeptide (I) comprising, to C direction, a region (R1) that recognises selectively a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAR-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                prodrug W24, containing, essentially, the single-chain Fv antibody OS4, specific for human fibroblast activation protein, trimerization linker,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 6; Page 47; 52pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYST-) UNIV STUTTGART (PFIZ/) PFIZENMAIER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunomodulatory and antiangiogenic activity, induce apoptosis and can be used for gene therapy. Kym-1 cells (2000) were incubated with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2002-362351/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunomodulatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wuest T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Moosmayer D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ħ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OS4; fibroblast activation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      treatment;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antianglogenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Grell M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nises selectively a specific component of the extracellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Scheurich P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tenascin;
                                                                                             fragment
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29 CAGAAÇACCGTGTGCACCGGATCCGCAGG 1 CAGAACACCGTGTGCACCTGCCATGCAGG

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41

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TTCCACCCGCACCTCCAGCTCCAC

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                            Matches
               Query Match
Best Local Similarity
                            Sequence 24
399 TTCCACCTTCACCTCCAGCTCCAC
            Conservative
                            BP; 4 A; 14 C; 2 G; 4 T; 0 other;
                3.6%;
91.7%;
            0;
                Score 20.8;
Pred. No. 1.
            Mismatches
                 1.8e+04;
                    DB 19;
            Indels
                    Length
                    24;
            0,
            Gaps
            0
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Query Match
Best Local Similarity
Matches 21; Conserv

399 TTCCACCTTCACCTCCAGCTCCAC 422

Conservative

0

3.3%;

Score 19.2; DB 19 Pred. No. 5.1e+04; Mismatches

DB 19;

Length Indels

the

0

Gaps

0;

Sequence 24 BP; 3 A; 14 C; 2 G; 5 T; 0 other;

Sequences shown in AAV55812 to AAV55827 represent primers used in the course of the invention for the multimerisation of minimal motifs. The course of the invention for the multimerisation of minimal motifs. The course of the invention for the multimerisation of minimal motifs. The course protein to proteolytic degradation that comprises linking or inserting conto or into the core protein a stabilising polypeptide of formula contour into the core protein a stabilising polypeptide of formula contour into the core protein a stabilising polypeptide of formula contour into the internative into a nucleic acid encoding a core protein. The fusion proteins of the invention are contour resistant to degradation by proteases and, thus, have a longer contour resistant to degradation by proteases and, thus, have a longer treating autoimmune diseases, cancer and inflammation. In particular, the core protein may be an IkappaB regulator protein for the treatment of inflammatory bowel disease, or a nitroreductase protein which can activate nitro drugs in enzyme/product therapy to treat cancer or other pathological conditions. The fusion proteins can also be used in course. And 9815 standard; DNA; Multiple Tiestion of minimal motifs using primer ZGS2 New fusion proteins resistant to proteolytic degradation comprising a core protein with a stabilising polypeptide WPI; 1998-312463/27 25-JUN-1997; 15-NOV-1996; Synthetic. Epstein-barr virus. 18-NOV-1998 AAV55815; Disclosure; Page 72; 120pp; English. 17-NOV-1997; 28-MAY-1998 WO9822577-A1 cancer; pathological condition; minimal Masucci MG (MASU/) MASUCCI M G. gustar protein; stabilising polypeptide; proteolytic degradation resistance; half-life; autoimmune disease; inflammation; nitro drug; likepas regulator protein; inflammatory bowel disease; in vivo imaging; nitroreductase protein; enzyme thorapy; prodrug therapy; protease; rancer; pathological condition; minimal motif; PCR primer; peptide sequence containing glycine repeats (first entry) 97US-0048945 96US-0030986 97WO-IB01508 24 comprising

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RESULT 7
AAV55817
ID AAV5
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ZZZZZZZZ
Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SX
                                                                                                                                                Sequences shown in AAV55812 to AAV55827 represent primers used in the course of the invention for the multimerisation of minimal motifs. The invention provides a method for increasing the resistance of a core protein to proteolytic degradation that comprises linking or inserting onto or into the core protein a stabilising polypeptide of formula ([Glya]X(Glyb)Y(Glyc)Z]n where Glya, Glyb, Glyc are 1-6 sequential Gly residues and X, Y, Z are Ala, Ser, Val, Ile, Leu, Met, Phe, Pro or Thr and n can be anything between 1-66. X, Y and Z need not be identical from n repeat to n repeat. Alternatively a nucleic acid encoding a stabilising polypeptide can be linked onto or inserted into a nucleic acid encoding a core protein. The fusion proteins of the invention are more resistant to degradation by proteases and, thus, have a longer transformance diseases. Cancer and inflammation. In particular, the products can be used for the invention are marked manner and the seases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New fusion proteins resistant to proteolytic degradation - comprising a core protein with a stabilising polypeptide comprising a peptide sequence containing glycine repeats
                                                    treating autoimmune diseases, cancer and inflammation. In particular, t core protein may be an IkappaB regulator protein for the treatment of inflammatory bowel disease, or a nitroreductase protein which can activate nitro drugs in enzyme/prodrug therapy to treat cancer or other pathological conditions. The fusion proteins can also be used in
                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 72; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Masucci MG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-JUN-1997;
15-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9822577-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Epstein-barr virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IkappaB regulator protein; inflammatory
nitroreductase protein; enzyme therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fusion protein; stabilising polypeptide; proteolytic degradation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Multimerisation of minimal motifs using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV55817;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAV55817 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-312463/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MASU/) MASUCCI M G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           resistance; half-life; auto:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gcer; pathologidal condition;
                                     methods such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       970S-0048945
960S-0030986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97WO-IB01508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24
                                     in vivo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disease;
                                 imaging.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bowel disease; in vivo imaging; produg therapy; professor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer
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ВP

AAX27663;

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RESULT 9
AAX27663
ID AAX2
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                                                                            Matches
       AAX27663 standard; DNA; 30
                                             26
                                                                                   Similarity
        ВÞ
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RESULT 8
AAF24737/c
ID AAF247
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                                                                                                                                                                                                                                                                                                                                  PCR primer used to amplify DNA encoding CDB-Tma peptide
                                                                                                                                                                                                                                                                                    Protest production; food processing; protein antibiotic; feed enzyme; CBD_rine; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF24737 Standard; DNA; 27
                                                                                                                                                                                                                                            United .
WPI; 2001-112219/12.
                               Shani Z,
                                                               (CBDT-) CBD TECHNOLOGIES LTD.
(YISS ) YISSUM RES DEV CO HEBREW UNIV JERUSALEM.
                                                                                                                                                    07-JUN-2000; 2000WO-IL00330.
                                                                                                                                                                                     21-DEC-2000.
                                                                                                                                                                                                                       WO200077174-'A1
                                                                                                                                                                                                                                                                                                                                                                         AFR. 2001
                                                                                                                  10-JUN-1999;
                             Shoseyov O;
                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                  9905-0329234.
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Expressing and isolating recombinant protein in a plant, useful for producing large quantities of recombinant proteins, by expressing a fusion protein including a cellulose binding peptide fused to a recombinant protein

Example; Page 48; 87pp; English.

The specification describes a method for expressing and isolating a recombinant protein in a plant. The method comprising expressing a fusion protein including the recombinant protein and a cellulose binding peptide fused to it, where the fusion protein is compartmentalised and sequestered within plant cells, plant derived tissue or cultured plant cells. The method is useful for obtaining large quantities of the recombinant proteins and protein products in a simple and cost-effective manner. Recombinant proteins may be used commercially, such as in the food processing industry, e.g. glucoamylases and glucose in healing processes, and to produce animal feed enzymes. PCR primers AAF24736-37 were used to amplify DNA encoding a CBD-Tma peptide. The amplified fragment was used to produce the fusion proteins of the such as in the food processing industry, e.g. glucoamylases and glucose isomerases are used for converting starch to high fructose corn syrup, proteinases for the hydrolysis of high molecular weight proteins and in manufacturing leather or alcoholic beverages, pectinesterases for pectin hydrolysis in food industry, lipases for cleaving ester linkage in triglycerides, and for effluent treatment. The recombinant proteins may further be used to produce protein antibiotics, which can be used

Sequence 27 BP; 7 ?

8888888888

Query Match Best Local S 520 TCCGACCCCATCCCCAACCCCCTT 543 3.3a; 87.5a; Score 19.2; DB 22 Pred. No. 5.3e+04; 0; Mismatches 3 DB 22; Length Indels 27; 0 Gaps

4 C; 12 G; 4 T; 0 other; 0 밁 ş SSXEREXEXE ABL51730 Claim 1; Page 5; 72pp; English

SEEKEKEK 20 JUL-1998; 21-JUL-1997; Synthetic gene; plant; gum; hydroxyproline-rich glycoprotein; HRGP; repetitive proline-rich protein; RFRP; arabino-galactan protein; AG Novel synthetic gene designed from repetitive peptide sequences hydroxyproline-rich glycoprotein WPI; 1999-132225/11. Kielszewski MJ Acacia sp. glycopeptide; DNA encoding a HRGP motif. 01-JUN-1999 (UYOH-) UNIV OHIO. 21-JUL-1998; 28-JAN-1999 WO9903978-A1 (first entry) 88. 980S-0897556. 970S-0897556. 98WO-US15083

AGP;

The invention relates to novel synthetic genes for plant gums. A new approach is described to the production of hydroxyproline-rich glycoproteins (HRGPs), repetitive proline-rich proteins (RRRPs) and arabino-galactan proteins (AGPs). Synthetic genes comprising a nucleic acid encoding the peptide (AAV01267) can be engineered for the production of repetitive glycopeptide modules in cells. The invention provided a new approach to the problem of producing plant gums that is not dependent on environmental factors and greatly simplifies the production of a variety of naturally occurring gums as well as designer gums. Note: The present nucleotide sequence is indicated as a peptide sequence in the claims.

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Sequence 30 BP; 6 A; 19 C; 0 <u>რ</u> G H 0 other;

Query Match Best Local : Matches Local . Similarity 21; Conserv Conservative 3.3%; ç Score 19.2; DB 20 Pred. No. 5.5e+04; D; Mismatches 3 DB 20; ü Indels Length 30; 0 Gaps

0

RESULT 10 ABL51730

ABL51730 standard; DNA; 30 ₽₽

09-JUL-2002 (first entry)

HRGP related oligonucleotide SEQ ID NO:10.

Plant; Gum arabic glycoprotein; HRGP; repétitive proline-rich pu AGP; plant gum; PCR primer; link protein; GAGP; hydroxyproline-rich glycoprotein; rotein; RPRP; arabinogalactan protein;

Acacia senegal Synthetic.

WO200178503-A2

12-APR-2000;

2000US-0547693. 2001WO-US12336.

12-APR-2001;

25-0CT-2001

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucles and proteins useful for producing hydroxy-proline rich
                                                                                                                                                                                                                                                                                     Hydroxyproline-rich glycoprotein (HRGP) related linker SEQ ID NO:38.
                                                                                                                                                                                                                                                                                                              09-JUL-2002
                                                                                                                                                                                                                                                                                                                                                           ABL51739 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present invention may be used to produce**HRGPs, repetitive proline-rich proteins (RPRPs) and arabinogalactan-proteins (AGPs) in plants via recombinant methodologies. Also described is the expression of synthetic genes designed from repetitive peptide sequences, such as glycoproteins (including the peptide sequences of gum arabic glycoprotein (GAGP)). ABL513730 to ABL51849 and ABS78401 to ABB78544 represent sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      other hydroxyproline (Hyp)-rich glycoproteins (HRGPs) and methods from the acids that encode them. The nucleic acids, proteins and methods from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kieliszewski MJ
         Nucleic acids and proteins useful
                                  WPI; 2002-041307/05
                                                                                                                                                                                                                                                            Plant; Gum arabic glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 4-5; 326pp;
                                                                                (UYOH-) UNIV OHIO
                                                                                                      12-APR-2000; 2000US-0547693
                                                                                                                              12-APR-2001; 2001WO-US12336
                                                                                                                                                                              WO200178503-A2
                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                               Acacia senegal.
                                                                                                                                                                                                                                      AGP; plant gum;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MIN OHIO
                                                                                                                                                                                                                                                                                                                                                                                                                                         401 CCACCTTCACCTCCAGCTCCACCT 424
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                                                                                                                                                                                                                                                                                                                                                                                                                    CCACCTTCACCTCCACCCCCATCT
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                                                                                                                                                                                                                                                  repetitive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6 A; 19 C; 0 G; 5 T; 0 other;
                                                                                                                                                                                                                                      PCR primer; li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               describes synthetic genes encoding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 19.2; I pred. No. 5.5e 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English
                                                                                                                                                                                                                                                                                                                                                                                                                    27
                                                                                                                                                                                                                                                  protein;
                                                                                                                                                                                                                                                GAGP; hydroxyproline-rich glycoprotein; rotein; RPRP; arabinogalactan protein;
           for
         producing hydroxy-proline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5.5e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               plant gums
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           rich
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glycoproteins in

plants

peptide

sequence

Disclosure; Page 72; 120pp; English.

New fusion proteins resistant to proteclytic degradation comprising a core protein with a stabilising polypeptide

comprising

glycine

WPI; Masucci

1998-312463/27

(MASU/) MASUCCI M G

25-JUN-1997; 15-NOV-1996; 17-NOV-1997;

97US-0048945 96US-0030986 97WO-IB01508 WO9822577-A1

Sequences shown in AAV55812 to AAV55827 represent primers used in the course of the invention for the multimerisation of minimal motifs. The invention provides a method for increasing the resistance of a core protein to proteolytic degradation that comprises linking or inserting onto or into the core protein a stabilising polypeptide of formula [(Glya)X(Glyb)Y(Glyc)Z]n where Glya, Glyb, Glyc are 1-6 sequential Gly residues and X, Y, Z are Ala, Ser, Val, Ile, Leu, Met, Phe, Pro or Thr

[(Glya)X(Glyb)Y(Glyc)Z] residues and X, Y, Z an and n can be anything k

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XXX2222222222XX
                                                                                                                                         The present invention describes synthetic genes encoding plant gums and other hydroxyproline (Hyp)-rich glycoproteins (HRGPs) and the nucleic acids that encode them. The nucleic acids, proteins and methods from the present invention may be used to produce HRGPs, repetitive proline-rich proteins (APRPs) and arabinogalactan-proteins (AGPs) in plants via
                                           'n
                                                                                                    recombinant methodologies. Also described is the expression of syntheti-
genes designed from repetitive peptide sequences, such as glycoproteins
                                                                                                                                                                                                                                                                       Example
                                                          including the peptide sequences of gum arabic glycoprotein (GAGP)).
                                         exemplification
                                                                                                                                                                                                                                                                       2; Page 53; 326pp; English
                                           of.
                                       the present
                                         invention
                                                                                                                          of synthetic
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                                Best Local
Matches 2
                                                   Query Match
                 401 CCACCTTCACCTCCAGCTCCACCT
4 CCACCTTCACCTCCACCCCCATCT
                                          Similarity
                                 Conservative
                                        3.3%;
                                  0
                                 Score 19.2; ] DB:
Pred. No. 5.5e+0.
0; Mismatches
                 424
27
                                           5e+04
                                                  24;
                                  Indels
                                                  Length
                                  0
                                  Gaps
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0

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Sequence 30 BP; 6 A; 19 C; 0 G; 5 T; 0 other;

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Synthetic
                                                                                                                                                        Fusion protein; stabilising polypeptide; proteolytic resistance; half-life; autoimmune disease; inflammat:
                                                                                                Epstein-barr
                                                                                                                            cancer; pathological
                                                                                                                                      IkappaB regulator protein; inflammatory
nitroreductase protein; enzyme therapy;
                                                                                                                                                                                    Multimerisation of minimal motifs using primer ZGY2.
                                                                                                                                                                                                         18-NOV-1998
                                                                                                                                                                                                                           AAV55821;
                                                                                                                                                                                                                                              AAV55821 standard; DNA; 24
                                                                                                                                                                                                        (first entry)
                                                                                                virus.
                                                                                                                            condition;
                                                                                                                                                                                                                                               ₽₽
                                                                                                                                               bowe]
                                                                                                                                                         inflammation;
                                                                                                                                                disease; in
                                                                                                                                                                   degradation;
                                                                                                                                               nitro drug;
n vivo imaging;
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stabilising

from n repeat to n repeat. Alternatively a nucleic acid encoding

polypeptide can be linked onto or inserted into a nucleic

acid encoding a core protein. The fusion proteins of the invention are more resistant to degradation by proteases and, thus, have a longer half-lifegithan the unfused core proteins. The products can be used for treating autoimmune diseases, cancer and inflammation. In particular, the core protein may be an IkappaB regulator protein for the treatment of activate intro drugs in enzyme/products protein which can activate intro drugs in enzyme/products therapy to treat cancer or other pathological conditions. The fusion proteins can also be used in

the

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14-AUG-1995;
16-SEP-1994;
16-SEP-1994;
          AAT17805-T17808, and AAT17810-T17813 represent glycosaminoglycan-degrading enzyme (GDE) inhibitors. The these sequences inhibit are endoglycosidases (which cleave
                                                                        Oligo:nucleotide(s) having sulphur substns. between nucleoside(s) for inhibiting glycosaminoglycan-degrading enzymes, for treating, e.g. cancer, inflammation, infection or autoimmune disorders.
glycosaminoglycan chains at
                                                     Claim 6; Page 33;
                                                                                                                     WPI; 1996-179936/18.
                                                                                                                                         Graham L,
                                                                                                                                                               (CARD-) CARDIAC CRC NOMINEES PTY
                                                                                                                                                                                                                                13-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                           cardiovascular
                                                                                                                                                                                                                                                                                                                                                                         disorder; infection; pathogenic or 
lar disease; vascular hyperplasia;
                                                                                                                                          Underwood
                                                                                                                                                                                     95AU-0004769.
94AU-0008226.
94AU-0008227.
                                                                                                                                                                                                                                95WO-AU00600
                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
1..30
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/note-
                                                    73pp; English.
                                                                                                                                          PA;
                                                                                                                                                                                                                                                                                               phosphorothicate,
internal sites), preferably heparanases (also
                                                                                                                                                                LID
                                                                                                                                                                                                                                                                                               or phosphorodithicate backbone"
                                                                                                                                                                                                                                                                                                                                                                           restenosis;
                      The GDEs which
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RESULT 14
ABL51740/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                   macrophages, neutrophils, leukocytes, endothelial cells, smooth muscle cells, carcinoma and tumour cells, and bacteria. They can also be used to inhibit smooth muscle cell activation, proliferation or migration. The sequences can be used to treat cancer, inflammation, autoimmune disorders, infection caused by pathogenic organisms, and cardiovascular disease, such as vascular hyperplasia, restenosis and atherosclerosis. These inhibitors can also be used as biochemical reagents for studying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    known as heparitinases) of mammalian or bacterial origin. These sequences can be used for inhibiting GDEs associated with platelets,
                                                             Hydroxyproline-rich glycoprotein (HRGP) related linker SEQ
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 30 BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                        GDE activities and mechanisms of enzyme activity.
HRGP; repetitive AGP; plant gum; F
                                Plant; Gum arabic glycoprotein;
                                                                                                 09-JUL-2002
                                                                                                                                    ABL51740
                                                                                                                                                                   ABL51740 standard;
                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                 (first entry
proline-rich
PCR primer; li
                                                                                                                                                                    DNA;
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Pred. No. 7.2e
0; Mismatches
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   in; GAGP; hydroxyproline-rich
h protein; RPRP; arabinogalact
linker; ss.
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                  arabinogalactan
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                                    glycoprotein;
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                  protein;
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TCCACCGCACCTCCAGCTCCA 23

807/c AAT17807 standard; DNA;

30

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Glycosaminoglycan-degrading enzyme; GDE; inhibitor; endoglycoheparanase; heparitinase; mammalian; bacterial; platelet; macheutrophil; leukocyte; endothelial cell; smooth muscle cell;

inhibitor; endoglycosidase;

macrophage;

tumour cell; activation; proliferation; migration; cancer; inflammation;

organism;

atherosclerosis;

Glycosaminoglycan-degrading enzyme inhibitor LG5PS.

30-OCT-1996 (first entry)

AAT17807;

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400 TCCACCTTCACCTCCAGCTCCA 421

Matches

20;

Conservative

, 0;

Mismatches

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Indels Length

1 24;

Gaps

0

bocar Similarity

90.98;

Score 18.8; DB 19 Pred. No. 6.6e+04; 4 T; 0 other;

DB 19;

Sequence 24 BP; 5 A; 13 C; 2 G;

diagnostic methods such as in vivo imaging.

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XXXXXXXX
present invention may be used to produce HRGPs, repetitive proline-rich proteins (RRPs) and arabinogalactan-proteins (AGPs) in plants via recombinant methodologies. Also described is the expression of synthetic genes designed from repetitive peptide sequences, such as glycoproteins (including the peptide sequences of gum arabic glycoprotein (GAGP)). ABL51730 to ABL51849 and ABPS01 to ABB78541 represent sequences used in the exemplification of the present invention.
                                                                                                                                                                                                               The present invention describes synthetic genes encoding plant gums and other hydroxyproline (Hyp)-rich glycoproteins (HRGPs) and the nucleic acids that encode them. The nucleic acids, proteins and methods from the
                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acids and proteins useful for producing hydroxy-proline rich glycoproteins in plants -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kieliszewski MJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-APR-2000; 2000US-0547693
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Acacia senegal.
                                                                                                                                                                                                                                                                                                                                            Example
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                                                                                                                                                                                                                                                                                                                                            2; Page 53; 326pp;
                                                                                                                                                                                                                                                                                                                                               English
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Query Match Best Local Similarity

3.2%; 0 C;

Score 18.4; DB 24; Pred. No. 9.4e+04;

Length 30

30 BP;

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Conservative

0,

Mismatches

0

Gaps

Search completed: May 21, 2003, 06:35:23 Job time: 223 secs

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comprises a method for validating a putative compound as a selective therapeutic compound towards a transcription factor response element. The method of the invention is useful for determining the transcriptional activation pathways used by any compound that is biologically active in a cell. This method allows a global view of gene transcription activation in response to diverse stimuli in multiple environments and is a significant improvement over case-by-case approaches, which would be limited to certain aspects of gene activation. It permits to save on clinical trials by screening properly the compounds that would have a lesser probability of providing undesirable, even severe side effects. The present sequence represents a double stranded oligonucleotide probe recognised by a specific transcription factor which is used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1993/c
1993/c
ABK97993 standard; DNA; 23 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A Cell-TRAP method, useful for producing or validating therapeutic compounds, by employing a recombinant cell-based library that carry constructs driven by a minimal promoter and a transcription factor-responsive element
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to a cell-TRAP method for selecting and producing a therapeutic compound which is presumed selective for, one or a restricted set of given transcriptional pathways and cell types by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 24; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-JUL-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Collimator method associated GATA mut oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK97993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comprising a reporter gene driven by a minimal promoter and a transcription factor responsive element (TFRE). The invention also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     restricted set of given transcriptional pathways and cell types by employing a recombinant cell-based library that carries a construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-575388/61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Blais Y, Rousseau P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-DEC-2000; 2000CA-2327581.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GENE-) GENEKA BIOTECHNOLOGY INC.
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23

457 CCCCGCAGAGAGGTGGCACCACC 479

Query Match
Best Local Similarity
Matches 20; Conserv

3.1%; Score 18.2; DB 24; 87.0%; Pred. No. 9.6e+04; ative 0; Mismatches 3;

Length 23; Indels

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Sequence 23 BP; 2 A; 9 C; 8 G; 4 T; 0 other;

5.1.4_p5_4578 Compugen Ltd.

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Minimum DB
Maximum DB
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No.
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Maximum Match 100%
Listing first 45 summaries
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seq length: 30
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Match
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
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   US-08-529-190B-10
US-08-629-190B-16
US-08-403-888A-34
US-08-192-102-15
US-08-192-861A-15
US-08-192-861A-15
US-09-106-038A-49
US-09-106-038A-50
US-09-106-038A-51
US-09-106-038A-52
US-09-106-038A-53
US-09-106-038A-54
US-09-106-038A-54
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2387.990 Million cell updates/sec
Sequence 19, Appl Sequence 7, Appl Sequence 7, Appl Sequence 15, Appl Sequence 16, Appl Sequence 16, Appl Sequence 17, Appl Sequence 18, Appl Sequence 15, Appl Sequence 15, Appl Sequence 15, Appl Sequence 16, Appl Sequence 17, Appl Sequence 48, Appl Sequence 50, Appl Sequence 51, Appl Sequence 51, Appl Sequence 51, Appl Sequence 52, Appl Sequence 53, Appl Sequence 54, Appl Sequence 55, Appl Sequence 56, Appl Sequence 57, Appl Sequence 58, Appl Sequence 59, Appl Sequence 51, A
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                                            Query Match
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Matches 21
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HYBRID PROTEINS 2 22 ESS: Y AND NEIMARK Th Street N.W., Ste. 300 ORM: PC-DOS/MS-DOS PC-DOS/MS-DOS IN Release #1.0, Version DATA: R: US/08/804,166 RR: US/08/804,166 RR: US/08/804,166 RR: COMPBELL-2A ROCKMATION: GRES: 5618 GRATION: GRES: 19: GRES: 19: STICS: palts id ngle	1 166-19/c ence 19, Application US/088 nt No. 6193972 reral INFORMATION: PPLICANT: Campbell, Robert PPLICANT: Clappel, Scott C	C 28 18 3.1 18 3 US-09-106-038A-62 C 29 18 3.1 18 3 US-09-106-038A-63 C 30 18 3.1 18 3 US-09-106-038A-63 C 31 18 3.1 18 3 US-09-106-038A-63 C 32 18 3.1 18 3 US-09-106-038A-63 C 33 18 3.1 18 3 US-09-106-038A-63 C 34 18 3.1 18 3 US-09-106-038A-63 C 35 18 3.1 18 3 US-09-106-038A-63 C 36 18 3.1 18 3 US-09-106-038A-63 C 37 18 3.1 18 3 US-09-106-038A-63 C 38 18 3.1 18 3 US-09-106-038A-63 C 38 18 3.1 18 4 US-09-136-139A-63 C 39 18 3.1 18 4 US-09-133-119-15 C 39 18 3.1 18 4 US-08-192-093A-15 C 39 18 3.1 18 4 US-08-192-093A-15 C 39 18 3.1 18 4 US-08-697-610-13 C 39 18 3.1 18 18 US-09-106-038A-63 C 30 18 3.1 18 3 US-09-106-038A-63 C 30 18 US-09-106-
ROTEINS N.W., Ste. N.W., Ste. 96 #1.0, Vers1 #04,166 996 996 996) »	-09-106-038A-62 -09-106-038A-63 -09-106-038A-65 -09-106-038A-65 -09-106-038A-66 -09-106-038A-68 -09-106-038A-69 -09-106-038A-69 -09-106-038A-69 -09-106-038A-69 -09-106-038A-69 -09-106-038A-69 -09-106-038A-70 -09-133-119-15 -08-192-093A-15 -08-192-093A-15 -08-192-093A-15 -08-474-542A-150 -08-477-542B-151 -08-457-648-151 -08-457-648-151
		Sequence

Similarity

Conservative

0

Score 21; Pred. No. Mismatches

DB 4; 10. 1.3e+03; 0;

Length 21; Indels

0

Gaps

0

142 ACTGAGGACTCAGGCACCACA 162

21 ACTGAGGACTCAGGCACCACA 1

Application US/08910991

Campbell;

MEDIUM TYPE: Floppy

Floppy disk

CITY: Washington STATE: D.C.

1: BROWDY AND NEIMARK
419 Seventh Street N.W., Ste. 300

HYBRID PROTEINS

COUNTRY:

USA

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

PatentIn Release #1.0,

Version #1.30

RRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/910,991 FILING DATE:

CLASSIFICATION: 530

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08-529-190B-7
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cent No. 583399
                                                                                                                                                                                                                          142 ACTGAGGACTCAGGCACCACA 162
                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RIOR APPLICATION DATA:
APPLICATION NUMBER: 08/804,166
FILING DATE: 20 February 1997
                                           ITLE OF INVENTION:
                                                                                                                                                                                               21
                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
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                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                          nucleic acid
                                                                                                                     Application US/08529190B
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Banner & Witcoff, Ltd
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                                       GLYCINE-CONTAINING SEQUENCES CONFERRING INVISIBILITY TO THE IMMUNE SYSTEM
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CLASSIFICATION:

GENT INFORMATION:

10-May-1993

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US-08-050-319B-15/c
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Best Local Similarity
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CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: SE9501
APPLICATION NUMBER: 1995
                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                ent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
               SOFTWARE: PatentIn Release #1.0, version #1.25 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                              399 TTCCACCTTCACCTCCAGCTCCAC 422
                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/529,190B FILING DATE: 15-SEP-1995 CLASSIFICATION: 514
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APPLICATION NUMBER: US/08/050,319B
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                                                                                                                                                Palo Alto
: California
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5633145
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                                                                                                                                                                            635 Bryant Street
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Pred. No. 1
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INFORMATION FOR SEO ID NO:

15:

CHARACTERISTICS: nucleic acid

30 base pairs

TELEFAX

(415

TELEPHONE: (415) 617-8999

REGISTRATION NUMBER:

5150-0030

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; MOLECULE TYPE:
US-08-465-982-15
                                      Query Match
Best Local Similarity
                                                                                                                                                                                 TELEFAX: (415) 327-33
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 'U
FILING DATE: 10-May-1
                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                           TELEPHONE: (415) 617-8999
                                                                                                        STRANDEDNESS:
TOPOLOGY: 11
                                                                                                                                                                                                                                                    NAME: Robbins, Roberta
REGISTRATION NUMBER: 3
13 CAGAACACCGTGTGCACCTG 32
                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER:
                                                                                                                                                     ENGTH:
                                                                                                                                     nucleic acid
                                                                                                                                                     30 base pairs
                                                                                                                                                                                              (415) 327-3231
                                                                                             DNA (genomic)
                                                                                                                        single
                                      3.4%;
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                          0
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                                       Score 20;
Pred. No
                                       3e+03;
                                                   Length 30;
                          Indels
                        Gaps
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Patent No.

GENERAL INFORMATION:

Masucci,

Maria G.
GLYCINE-CONTAINING SEQUENCES
CONFERRING INVISIBILITY TO THE IMMUNE SYSTEM

0

CORRESPONDENCE

ADDRESS:

ADDRESSEE:

STREET:

One Financial Center

Banner & Witcoff, Ltd

Boston

NUMBER OF SEQUENCES:

ITLE OF INVENTION:

SOFTWARE: PatentIn Rel

Patentin Release #1.0,

APPLICATION NUMBER:

US/08/465,982

CLASSIFICATION: FILING DATE OMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk

94301

STREET:

Palo Alto

635 Bryant Street

Reed & Robbins

California

USA

RRESPONDENCE ADDRESS:

DDRESSEE:

COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS

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Query Match
Best Local Similarity
Watches 21; Conserva
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                          ·08-529-190B-16
                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
                                                                                                            399 TTCCACCTTCACCTCCAGCTCCAC 422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30 CAGAACACCGTGTGCACCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: DOS
SOFTWARE: Wordperfect 6.1
                                                                                  1 TTCCACCCGCACCTCCAGCTCCTC 24
                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                               POPOLOGY:
                                                                                                                                                                                                                                                                                                               TELEPHONE:
                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DDRESSEE:
                                                                                                                                                                                                                                                       ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OUNTRY:
16, Application US/08529190B
o. 5833991
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                                                                                                                                                                                                                                     nucleic acid
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                                                                                                                                        Conservative
                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IBM Compatible
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                                                                                                                                                                                                                           single
                                                                                                                                                                                                                                                                                                                                                                                                     01-SEP-1995
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                                                                                                                                                       3.3%;
87.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GLYCINE-CONTAINING SEQUENCES
                                                                                                                                                                                                                                                                                                                                                                                                                 US08/522,595
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                                                                                                                                                                                                                                                                                                                                    34,380
NER: 3255/53015
                                                                                                                                                                                                                                                                                                                                                                   Kathleen A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11
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                                                                                                                                                       Score 19.2;
Pred. No. 4
                                                                                                                                                                 2; Length 24;
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13 CAGAACACCGTGTGCACCTG 32

Similarity

100.08;

Score 20; Pred. No

DB 1; 3e+03;

Length 30;

Indels.

Ç.4.

0,

Conservative

. 0;

TYPE: DNA (genomic)

linear

single

30 CAGAACACCGTGTGCACCTG 11

US-08-465-982-15/c

ent No.

Application US/08465982

INFORM?

ITLE OF INVENTION: Modified human INFalpha (Tumor

Modified

M.J.C. Turner, Feldmann,

F.M Brennar

OMPUTER READABLE FORM

EDIUM TYPE:

IBM Compatible

Diskette

CATION NUMBER:

US/08/529,190B

15-SEP-1995

ING SYSTEM: DOS

RE: Wordperfect 6.1

REPLICATION DATA:

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)8-403-888A-33/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ent No.
                                                                                                               SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                  COMPUTER: READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 1.44 Mb
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   400 TCCACCTTCACCTCCAGCTCCA 421
APPLICATION NUMBER: US/08/403,888A FILING DATE: 12-UN-1995 CLASSIFICATION: 435 TOR APPLICATION DATA: APPLICATION NUMBER: 07/954,185 FILING DATE: 29-SEP-1992
                                                                                                                                                                                                                                                                                                                                                                                                  PPLICANT: Hanecak et al
ITLE OF INVENTION: Olig
ITLE OF INVENTION: Sequ
                                                                                                                                      OPERATING SYSTEM:
SOFTWARE: Wordper
                                                                                                                                                                                                                                                                                                                                                       RRESPONDENCE ADDRESS
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5. 5952490
                                                                                                                                                                                                                                       19103
                                                                                                                                                                                                                                                                                              Philadelphia
                                                                                                                                                                                                                                                                                                                3: Woodcock Washburn Kurtz Mackiewicz One Liberty Place - 46th Floor
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RESULT 9
US-08-403-888A-34/c
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                                                                                    US-08-403-888A-34
                                          Query Match
Best Local 9
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                            Matches
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                                                                                                                                                                           INFORMATION FOR SEQ
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                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: 3.5 inc
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                                                                                                                                                                                                                                                                           FILING DATE: 29-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           518 CCTCCGACCCCATCCCCAACCCC 540
                                                                                              TYPE: nucleic acid
STRANDEDNESS: singl
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                               REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/403,888A FILING DATE: 12-JUN-1995
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OR APPLICATION DATA:
07/954,185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LE OF INVENTION:
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CCTCCGACCCCATCCCCAACCCC 540
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                                          Similarity
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One Liberty Place - 4
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                                           Score 18.2;
Pred. No. 9
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Pred. No. 9
                            Mismatches
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                                                      Length 25;
                            Indels
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Query Match

Local

Similarity

3.2%;

Score 18.8; Pred. No. 6

B

Length 24;

0

COLECULE TYPE:

OLOGY:

linear

other nucleic acid

STRANDEDNESS:

nucleic acid

TELEFAX: 617-345-9111 NFORMATION FOR SEQ ID NO:

ELEPHONE:

NAME: Williams, Ph.D., Kathleen A REGISTRATION NUMBER: 34,380 REFERENCE/DOCKET NUMBER: 3255/53015

MUNICATION INFORMATION:

ORNEY/AGENT INFORMATION:

TATION NUMBER:

ON NUMBER: CATION DATA:

SE9501324-9

10-APR-1995

ICATION DATA; US08/522,595

01-SEP-1995

SEQUENCE CHARACTERISTICS:

15, Application US/08192102

Junming

1, Peter E.

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                                                                                     MOLECULE TYPE:
                                                               Query Match
           109 TTGTGCCTACCCCAGATT 126
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18
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                                                                                                            OPOLOGY:
                                                     Similarity
                                                                                                                                 nucleic acid
                                                                                                                                                                                                                                                                                                  ATION NUMBER:
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                                                                                                                                                                                                                                                                   ION NUMBER:
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18-MAR-1991
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29-JAN-1993
                                                                                                                                                                                                                                                                                      18-MAR-1992
                                                                                                                                                                                                                                                                                                                      MBER: US 07/943,852
11-SEP-1992
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                                                                                                                                                                                                                     22,592
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                                                   DB 1; L
9.4e+03;
                                                               Length 18;
                                         Indels
                                         0;
                                         Gaps
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NEORMATION FOR SEQ
                                                                      SEQUENCE CHARACTERISTICS
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                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
OR APPLICATION DATA:
OB/192,093
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/0
FILING DATE: 18-OCT-1994
                          STRANDEDNESS:
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                                                                                                                                                         REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                               OR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OMPUTER: IBM PC compatible
PERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DIUM TYPE: Floppy
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                                                                                                                                                                                                              CATION NUMBER:
TYPE: CDNA
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                                                        18 base pairs
                                                                                                                                                                                                                                                       ON NUMBER:
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                                                                                                                                          DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER:
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                                                                                                                                                                                                                                    UMBER: US 07/853,606
18-MAR-1992
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                                                                                                                                                                                                  IMBER: US 07/670,827
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                                                                                                                                                                      David E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-FEB-1994
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                                                                                                                                                                                                                                                                                                                             MBER: US 08/010,406
29-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                      MBER: US 08/013,413
02-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                                              04-FEB-1994
                          single
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                                                                                    ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peter E
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                                                                                                                                            NYU93-01M4
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

Lexington Massachusetts

USA

: Hamilton, Brook, Smith & Reynolds, P.C. Two Militia Drive

ANTIBODIES AND ASSAYS EMPLOYING ANTIBODIES

APPLICATION NUMBER: US,

SOFTWARE:

Patentin Release #1.0, Version #1.25

PILING DATE: 04-FEB-193.
PILING DATE: 04-FEB-193.
APPLICATION NUMBER: US 08/013,413
PILING DATE: 02-FEB-1993

ASSIFICATION: 424
R APPLICATION DATA:
PLICATION NUMBER: US/08/192,093

04-FEB-1994

US/08/192,102

Query Match 3.3 Best Local Similarity 100 Matches 18; Conservative

Length 18 Indels

0

Gaps

0

RESULT 11

18 TTGTGCCTACCCCAGATT 1

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US-08-192-861A-15
                                                         Matches
                                                                       Best Local
                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8-192-861A
                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
                                                                                                                                                                                                                                             TELEPHONE: (781) 861-6240
                                                                                                                                 MOLECULE TYPE:
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MEDLIM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
                           109 TIGTGCCTACCCCAGATT 126
                                                                                                                                                                                                                                                                                                                    TTORNEY/AGENT INFORMATION:
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PRICATION DATA:
US 08/013,413
18 TIGIGCCTACCCCAGATI 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Lexington
STATE: Massachusetts
                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 29-JA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/192,861A FILING DATE: 04-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                      NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
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                                                                                                                                                                                                                                                                                                                                                              APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                               CATION NUMBER: US 07/853,606 NG DATE: 18-MAR-1992
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                                                                       Similarity
                                                                                                                                                                    nucleic acid
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ON NUMBER: US 07/943,852
NTE: 11-SEP-1992
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                                                         Conservative
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29-JAN-1993
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                                                                     Score 18; DB 2; 
; Pred. No. 9.4e+03
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                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                     quence 48, Application US/09106038A tent No. 6007995
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                                    CURRENT APPLICATION DATA
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                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 1.44 Mb
                                                                                                                                                                      STREET: 22.
                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: Windows NT SOFTWARE: Microsoft Word 97
                                                                                                                                                                                                                                                                         TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                            18 GGAGAAACAGAACACCGT 1
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NAME: Laurel Spear Bernstein
REGISTRATION NUMBER: 37,280
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TATE: CA
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LING DATE: June 26, 1998
ASSIFICATION: 514
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VENTION: ANTISENSE MODULATION OF TNFR1
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NUMBER: US/09/106,038A
: June 26, 1998
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REFERENCE/DOCKET NUMBER: ROTELECOMMUNICATION INFORMATION

RTS-0004

760)

PELEPTE: (760) 603-3820 RMATION FOR SEQ ID NO: QUENCE CHARACTERISTICS:

DNESS: nucleic acid

single

linear

CLASSIFICATION: 514
ATTORNEY, AGENT INFORMATION:
NAME: Laurel Spear Bernstein
REGISTRATION NUMBER: 37,280

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18 CGAGTGTGTCTCCTGTAG 1

Query Match
Best Local Similarity
Matches 18; Conserva

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Mismatches

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Best Local Similarity 100.0%;
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                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Laurel Spear Bernstein
REGISTRATION NUMBER: 37,280
REFERENCE/DOCKET NUMBER: RTS-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (760) 931-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          equence 49, Application US/09106038A stent No. 6007995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Brenda
                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 1.44 Mb
COMPUTER: IBM PC compatible
OCMEDIUM SYSTEM: Windows NT
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
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COUNTRY: U.
ZIP: 92008
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/106,038A FILING DATE: June 26, 1998 CLASSIFICATION: 514
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TENTION: ANTISENSE MODULATION OF THER1
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Search completed: May 21, 2003, 07:31:30 Job time: 77 secs

GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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                                     Sequence 128, App
Sequence 15, Appl
Sequence 11, Appl
Sequence 21, Appl
Sequence 21, Appl
Sequence 221, App
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Sequence 9, Appl1
Sequence 9, Appl1
Sequence 19, Appl
Sequence 7, Appl
Sequence 128, App
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Patent No. US20020090676A1
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                                                                                                                                                                                                                                APPLICANT: Hauptmann, Rudolph
APPLICANT: Himmler, Adolph
APPLICANT: Maurer-Fogy, Ingrid
APPLICANT: Stratowa, Christian
APPLICANT: The Receptors, TNF Binding Proteins and DNAs Coding for INVENTION: Them
                                  RIOR APPLICATION NUMBER: 08/383,676
RIOR FILING DATE: 1995-02-01
RIOR APPLICATION NUMBER: 08/153,287
RIOR FILING DATE: 1993-11-17
RIOR APPLICATION NUMBER: 07/821,750
RIOR FILING DATE: 1992-01-02
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                                                                                                                                                    JERENT APPLICATION NUMBER: US/09/899,422
JERENT FILING DATE: 2001-08-21
RIOR APPLICATION NUMBER: 09/525,998
RIOR FILING DATE: 2000-03-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: CDS
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                APPLICATION NUMBER: 07/511,430
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LING DATE: 1993-11-17
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LING DATE: 1995-02-01
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Pred. No.
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; NAME/KEY: CDS

; LOCATION: (1)..(30)

US-09-899-422-9
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                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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                        142 ACTGAGGACTCAGGCACCACA 162
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                                                            Local
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ZIP: 22207
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                                                                                                                                                                  ENGTH:
                                                                                                                ECULE TYPE:
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E: Browdy, Roger L.
SISTRATION NUMBER: 25,
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                                                           Similarity 100.0%;
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                                                                                                                                                                  21 base pairs
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Patent No.

US20020064771A1

PPLICANT: Zhong, Weldong

INFORMATION:

CANT: Ferrari, Eric

URRENT APPLICATION NUMBER: US/09/828,034
URRENT FILING DATE: 2001-04-06
PRIOR ARPLICATION NUMBER: U.S. 60/195,852
PRIOR ARELICATION SUMBER: 2000-04-06
UNBER 40 40 10 NOS: 33

atentIn Ver. 2.1

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RESULT 7
US-10-043-432-15/c
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; Publication No. U
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S-10-113-877-128/c
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Best Local Similarity /0...
23; Conservative
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                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version 4.0 EQ ID NO 128 LENGTH: 23
                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                 457 CCCCGCAGAGAGGTGGCACCACC 479
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US20020177218A1
                                                                                                                                                                                                                                                                                                                                                                                      CATION NUMBER: 60/314,330
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   Application US/10043432
10. US20030054004A1
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                                                                                                                                                          3.1%;
ilarity 87.0%;
Conservative
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Pred. No. 2
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Sequence 15, Application US/09756301A Patent No. US20010027249A1

09-756-301A-15/c

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GENERAL INFORMATION:

INVENTION: Anti-TNF Antibodies INVENTION: Human Tumor Necros

and Peptides Factor

Human Tumor

Daddona, Peter Ghrayeb, John

Knight, David M.

Scott

Vilcek, Jan

Junming

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                                                                                                              ) OTHER INFORMATION: PCR oligonucleotides US-10-043-432-15
                                                                                                                                                      NUMBER OF SEQ ID NOS: 19
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 15
LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
                                                        Matches
                                                                    Query Match
Best Local (
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              109 TTGTGCCTACCCCAGATT 126
18 TTGTGCCTACCCCAGATT 1
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LING DATE: 1994-1
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                                                                       Similarity
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                                                        Conservative
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Pred. No.
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                                                                                                                                                                                                                                                          07/670,827
                                                                                   Length 18
                                                       Indels
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JS-09-828-034-7

OTHER INFORMATION: Description of Artificial Sequence:

3.2%;

Score 18.8; D Pred. No. 1.6e 0; Mismatches

.6e+04;

DB 10;

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Synthetic

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ORGANISM: Artificial Sequence

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Query Match
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                                                                                                                                                                                                                                                                               PPLICANY: Siegel, Scott
TTLE OF INVENTION: Anti-TANF AntiDodies and Peptides of
ITLE OF INVENTION: Human Tumor Necrosis Factor
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                                                                                                                                                                                    APPLICATION NUMBER: U.
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                                                                          APPLICATION NUMBER: U.S
FILING DATE: 1994-02-04
                                                                                                        APPLICATION NUMBER: U.S. FILING DATE: 1994-10-18
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                                                             APPLICATION NUMBER: U.
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LING DATE: 1994-02-04
PLICATION NUMBER: U.S.
LING DATE: 1994-02-04
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DATE: 1994-02-0
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S. 08/010,406
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Pred. No.
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                                                            SOFTWARE: FastSEQ
SEQ ID NO 15
LENGTH: 18
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Best Local Similarity
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                                                                                                                        PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                 URRENT FILING DATE: 2001-01
RIOR APPLICATION NUMBER: U.S
RIOR FILING DATE: 1998-08-12
RIOR APPLICATION NUMBER: U.S
RIOR ETLING DATE: 1995-12-11
OTHER INFORMATION: PCR oligonucleotides
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                              )RGANISM: Artificial Sequence
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LING DATE: 1993-02-02
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                                                                                                                                                                                                                                                                              ING DATE: 1994-02-
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RIOR FILING DATE: 1995-12-1
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RIOR FILING DATE: 1998-08-1
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ITLE OF INVENTION: Anti-TNF Antibodies and Peptides
ITLE OF INVENTION: Human Tumor Necrosis Factor
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CURRENT FILING DATE: 2001-12-07
PRIOR APPLICATION NUMBER: US/09/927,703
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TANGE: DNA
ORGANISM: Artificial Sequence
FEATURE:
FEATURE:
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OTHER INFORMATION: PCR oligonucleotides US-10-043-450-15
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TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: PCR oligonucleotides
10-044-534-15
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NUMBER OF SEQ ID NOS: 19
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CATION NUMBER: U.S. 07/670,827
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ATION FOR SEQ ID NO:
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R APPLICATION DATA:
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AL455550 T. bruce1
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ALIGNMENTS

										
		JOURNAL COMMENT	TITLE		REFERENCE AUTHORS		ORGANISM	KEYWORDS	ACCESSION	RESULT 1 AZ776616 LOCUS DEFINITION
Hall2, USA Tel: 801 585 5606 Tex: 801 585 7177 Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Error: 0.00	University of Utah University of Utah University of Utah Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT	plasmid inserts Unpublished (2000) Contact: Robert B. Weiss	and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb	<pre>Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Rellly ,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.</pre>	<pre>l (bases 1 to 28) Dunn,D., Aoyag1,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,</pre>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.	nouse mouse. Mus musculus	GSS.	clone UUGC2M0010K24 F, DNA sequence. AZ776616 AZ776616 GT:120A394	A2776616 28 bp DNA linear GSS 16-FEB-2001 2M0010K24F Mouse 10kb plasmid UUGC1M library Mus musculus genomic

Seq primer: CGTTGTAAAACGACGGCCAGTClass: plasmid ends

column:

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REFERENCE
AUTHORS
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TA42E02P/c
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                                                                                            Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically shear
                                                                                                                                                                       Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucel genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, HiCambridge CB10 15A, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                    Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Trypanosoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Trypanosoma brucei.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AL455550.1 GI:11856678
                                described in detail
                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
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                 give a tight size distribution (b). The v + 1 method used for the library construction is scribed in detail in Smith, H. and Venter, J.C. (Making small)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymether and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           musculus C57BL/6J
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75.0%;
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whole genome shotgun sequencing projects.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 16.8; DB 17;
Pred. No: 6.9e+06;
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RESULT 4 AZ788326/c

DEFINITION

AZ788326

2M0035P16F Mouse 10kb plasmid $\overline{\text{UUGC1M}}$ library Mus musculus clone $\overline{\text{UUGC2M0035P16}}$ F, DNA sequence.

GSS 16-FEB-2001

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BASE COUNT
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CCACCGGTGGAGCTCCAAGCTTT 4
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920 E. 58th Street, Chicago,
                                                                                                                                                                                                                                                                                                                                                      Contact: Turkewitz AP
Molecular Genetics and Cell Biology
                                                                                                                                                                                                                                                                                                                                                                                     ,J. and Kloburcher,L.
EST from Tetrahymena thermophila, strain CU428.1, growing cells
                                                                                                                                                                                                                                                                                                                                                                                                                                              Tetranymena thermophila
Tetrahymena thermophila
Eukaryota; Alveolata; Cillophora; Oligohymenophorea;
Eukaryota; Alveolata; Tetrahymena.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5009-0-62-A04.t.1 Chilcoat/Turkewitz cDNA (large fraction)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Barrell, Oxford Universionall: nelsayed@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tetrahymena thermophila.
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primer: T3.
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                                                                                                                          /note-"Vector: BlueScript2 SK+; Details on library preparation can be found in Chilcoat and Turkewitz (2001) Proc. Natl. Acad. Sci USA, 98: 8709-8713."
7 c 8 g 4 t
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                                                                                                                                                         /organism="Tetrahymena thermophila"
/strain="CU428.1"
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/strain-"TREU927"
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A2464926 30 bp DNA linear 1M0274J04R Mouse 10kb plasmid UUGC1M library Mus clone UUGC1M0274J04 R, DNA sequence.
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Mammalla; Eutherla; Rodentia; Sciurognathi; Muridae; Murinae; Mus
(bases 1 to 19)
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Insert Length: 10000 Std Erro
Plate: 0035 row: P column: 10
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                         chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance. The following formula 6\ c 3\ g 3\ t
                                                                                                                                                                                                                                                                                                                                                                                                                                                               adaptored vector DNA, and transformed into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   with adaptors complementary to the insert adaptors purified. The sheared, adaptored mouse DNA was annu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were bligated to the blunt ends in high molar excess. The daptored DNA was purified and size-selected for a 9.5 to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (http://www.jax.org/resources/documents/dnares/). The DNF was hydrodynamically sheared by repeated passage through DNA inch orifice at constant velocity. The sheared DNA
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/strain="C57BL/6J"
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'clone="UUGC2M0035P16"
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note="Vector: PWD42nv; Purified genomic DNA from M.
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                                                                                                                                                                                                                                                                                            2.8%;
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                                                                                                                                                                                                                                                                     Score 16.4; DB 17;
Pred. No. 8.3e+06;
0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.
Insert Length: 10000 Std
Plate: 0274 row: J column
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Fax: 801 585 7177
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Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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Location/Qualifiers
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                                                                                                                                                                                                    10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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note="Vector: PWD42nv; Purified genomic DNA from M.
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AZ875577 30 bp DNA 11 2M0190G06F Mouse 10kb plasmid UUGC1M library clone UUGC2M0190G06 F, DNA sequence.

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BASE COUNT
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AA936737 25 bp mRNA o159f10.sl NCI_CGAP_HN4 Homo sapiens cDNA similar to TR:Q18444 Q18444 COSMID C34D4.
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Insert Length: 10000 Std Error: 0.00
Plate: 0190 row: G column: 06
Seg primer: CGTTGTAAAACGACGGCCAGT
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Fax: 801 585 7177
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University of Utah
Rm. 308, Biomedical Polymers Res
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stiand selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Mus musculus"
strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ocation/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sculus C57BL/6J (male) was obtained
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        lone_lib="Mouse 10kb plasmid UUGCIM library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 _xref="taxon:10090"
one="UUGC2M0190G06"
                                                                                                                                                                                                                                                                                                   2.8%;
72.4%;
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--"Vector: PWD42nv; Purified genomic DNA from M.
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                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                                                Score 16.2; DB 17 Pred. No. 9.7e+06;
                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                            29
                                                                                                                                                                                                                                                                                                                                                                                                                            coli XL10-Gold (Stratagene) cells
                                                                                                                                                                                                                                                                                                                          DB 17;
                    RNA linear EST 29-APR-1998 CDNA clone IMAGE:1486987 3'
     ;, mRNA sequence
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                                                                                                                                                                                                                                                                                                                       Length 30;
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JOURNAL COMMENT
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VERSION
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AZ348233
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                                                                                                                                                                                                                                                                                                                                                                   AZ348233 25 bp DNA linear GSS 29-SEP-200 1M0084G04R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0084G04 R, DNA sequence.
University of Utah
                                             plasmid inserts
Unpublished (2000)
                                                                                     ,M., Rose,M., Rose,R.,
and Wright,D.,Weiss,R.
Mouse whole genome scal
                                                                                                                              Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C. Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Re,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhauser
                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 25)
                                                                                                                                                                                                                                                               Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
Tissue Procurement: John Ensley, M.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CDNA Library Preparation: Stratagene, Inc. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequebring by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 25)

NCI/NIDR-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Dental Research,

Cancer Genome Anatomy Project (CGAP), Tumor Gene Index

Unpublished (1997)
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AA936737.1 GI:3094771
                                                                                                                                                                                                                                                                                        house mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ontact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                           Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligo dT. Average insert size 1.5 kb. 5' adaptor sequence: 5' GAATTCGGCACGAG 3' 3' adaptor sequence: 5' (GA )10ACTAGTCTCGAGTTTTTTTTTTTTTTTT 3'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note-"Organ: pharynx; Vector: Bluescript SK-; Site_1: ECCRI; Site_2: XhoI; Cloned unidirectionally. Primer:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /tissue_type="squamous cell carcinoma"
/lab_host="SOLR (kanamycin resistant)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone="IMAGE:1486987"
/clone_11b="NCI_CGAP_HN4"
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       79.28;
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0;
    Genome Center
                           Weiss
                                                                                   scaffolding with paired end reads
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Pred. No.
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5;
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KEYWORDS
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AZ823359
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 26)
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Injert Length: 10000 Std Error: (
Plate: 0084 row: G column: 04
Seq primer: CACACAGGAAACAGCTATGACC
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Fax: 801 585 7177
University of Utah University of Utah
                                                               Unpublished (2000)
                                                                                          plasmid
                                                                                                         and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                 Dunn,D., Aoyag1,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
                                                                                                                                                                                                                                                                                                                                                                                                                          2M0097C19F Mouse 10kb plasmid UUGC1N clone UUGC2M0097C19 F, DNA sequence
                                            Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                              AZ823359.1 GI:12993267
                                                                                                                                                                                                                                                                                                                                  nouse mouse.
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quality sequence stop: 25.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (http://www.jax.org/resources/documents/dhares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /lab_host="E. Coli strain XL10-Gold, T1-registant, F-"
/note="Vector: PWD42nv; Purified genomic AMA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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79.28;
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                                                                                                                                                          Rose, R., Stokes, R., Tingey, A., von
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Pred. No.
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smid UUGCIM library Mus musculus
                      Center
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. 1.1e+07;
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RESULT 10 AZ343341

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ACCCCATTCAGGAGTGGGGCCACA 26

536 ACCCCCTTCAGAAGTGGGAGGACA 559

COMMENT

JOURNAL TITLE

Inpublished plasmid inserts

(2000)

Robert B.

Utah Utah

Genome Center Weiss REFERENCE

AUTHORS

Dunn, D., Aoyagi, A., Mammalia; Eutheria; 1 (bases 1 to 27)

and Wright, D., Weiss, R.

whole genome

scaffolding with paired end reads from 10kk

Slam, H., Longacre, S., Mahmoud, M., Meenen, E., M., Rose, M., Rose, R., Stokes, R., Tingey, A.,

Barber, M., Beacorn, T.,

Meenen, E., Pedersen, T.,

Duval, B., Hamil, C.,

SOURCE KEYWORDS VERSION ACCESSION DEFINITION SD209

SS

AZ343341

10kb plasmid

27 bp DNA musculus

29-SEP-2000

genomic

DNA sequence.

AZ343341.1 GI:10421314 1M0076B04R Mouse 10kb pclone UUGC1M0076B04 R,

ORGANISM

Mus musculus

house mouse.

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus

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BASE COUNT
ORIGIN
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  Matches
                           Query Match
Best Local
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                         Local
     19;
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High quality sequence stop: 26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erry
Plate: 0097 row: C column: 1
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Fax: 801 585 7177
                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seq primer: CGTTGTAAAACGACGGCCAGT
  Conservative
                                                                                                                                                                                                                                         ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWn942 (gil4732114|gb)AF129072.1), a copy number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biomedical Polymers Research Bldg., 20 S.
                                                                                                                    adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

8 c 9 g 3 t
                                                                                                                                                                                                                                                                                                                                                                                       (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                          purified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 musculus C57BL/6J (male) was obtained
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="Vector:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="UUGC2M0097C19"
/clone_11b="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'strain-"C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          aboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lab_host="E. Coli strain XL10-Gold, T1-resistant, F
note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  organism-"Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ocation/Qualifiers
                         2.7%;
                                                                                                                                                                                                                     The sheared, adaptored mouse DNA was annealed to
0;
                         Score 16; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Std Error:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                musculus
     Mismatches
                                                  DB 17;
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                           .1e+07;
                                             Length 26;
  Indels
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  Gaps
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308, Biomedical Polymers Research Bldg.,

20 S.

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BASE COUNT
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ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                                                                       DEFINITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 11
                                                                                                            TITLE
                                                                                                                                                                                                   AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 254 AGCTCTACTCCATTGTTTGTGGGA 277
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AACACTCCTCCATTGCTGGTGGGA 26
                                                                                                  University of Utah Genome Center University of Utah
                                                              Unpublished (2000)
                                                                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                           1M0290M03R Mouse 10kb plasmid \bar{U}UGC1M library Mus musculus clone \bar{U}UGC1M0290M03 R, DNA sequence.
                                           Contact: Robert B.
                                                                                  plasmid inserts
                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                      AZ474207.1 GI:10632332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High quality sequence stop: 27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Ingert Length: 10000 Std Erro
Elate: 0076 row: B column: 0
                                                                                                                                                                                                                                                                                                             nouse mouse
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801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
7 c 8 g 7 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          as hydrodynamically sheared by repeated passage through .005 inch orifice at constant velocity. The sheared DNA as blunt end repaired with T4 DNA polymerase and T4 Olynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  _xref="taxon:10090"
lone="UUGC1M0076B04"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         host="E. Coli strain XL10-Gold, T1-registant, F
|-- "Vector: PWD42nv; Purified genomic_part from M.
                                       Weiss
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Pred. No.
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1.1e+07;
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                                                                                                                                                                                                                                          Murinae; Mus.
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Contact: Robert B. Weiss University of Utah Genome Center University of Utah

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JOURNAL
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AUTHORS
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VERSION
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AZ994586
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Best Local 9
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                                                                                                    Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                  Unpublished (2000)
                                    plasmid inserts
                                                             Mouse whole genome
                                                                               and Wright, D., Weiss, R.
                                                                                                                                                                      Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
1 (bases 1 to 26)
                                                                                                                                                                                                                                        Mus musculus
                                                                                                                                                                                                                                                                                                       AZ994586.1
                                                                                                                                                                                                                                                                                                                                                clone UUGC2M0280B08
                                                                                                                                                                                                                                                                                                                                                                     2M0280B08F, Mouse
                                                                                                                                                                                                                                                                 nouse mouse.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.
Insert Length: 10000 Std
Plate: 0290 row: M column
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 adaptored vector DNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DN
was hydrodynamically sheared by repeated passage through
0.005 inch orifice at constant velocity. The sheared DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      one_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                                              10kb plasmid UUGC2M
108 F, DNA sequence.
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                                                           scaffolding with paired end reads from 10kb
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Pred. No.
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mid UUGC2M library Mus musculus genomic
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                                                                                                                                                                                            Craniata; Vertebrata;
Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 17;
1.1e+07;
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; Murinae; Mus
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Best Local
                                                                                                                                                                                                   AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                     1M0095G10R Mouse 10kb plasmid clone UUGC1M0095G10 R, DNA sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
17; Conserv
                                                            plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 27)
                                                                                                                                                                                                                                                                                          Mus musculus
University of Utah
University of Utah
                                                                                                                             ,M., ROSe,M., ROSe,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Re.
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                                                                                                      Mouse whole genome scaffolding with paired end reads from 10kb
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                               nouse mouse.
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quality sequence stop:
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                                     Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative electrophoresis. Vector DNA was prepared from a derivative of pWD42 (g114732114) gb laFT129072.1), a copy-number inducible derivative of plasmid RL. The vector was ligated with adaptors complementary to the insert adaptors and
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/note="Vector: PWD42nv; Purified genomic with from M.
musculus C57BL/6J (female) was obtained from the Jackson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       organism="Mus musculus"
strain="C57BL/6J"
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'clone="UUGC2M0280B08"
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                                                                                                                                                                                                                                                                                                                                                           GI:10468500
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89.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     lone_lib="Mouse 10kb plasmid UUGC2M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          selected for ampicillin resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                :p://www.jax.org/resources/documents/dnares/).
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                    Genome Center
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                                         Weiss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.8; DB 17
Pred. No. 1.2e+07;
                                                                                                                                                                                                                                                                                                                                                                                                     DNA sequence.
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E. coli XL10-Gold (Stratagene) cells
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                                                                                                              Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reill)
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                           SSD
                                                                                                                                                                                                                                                                                                                                                                     AZ842796
                                                                                                                                                                                                                                                                                                                                                                                      2M0141120F Mouse 10kb plasmid TUGC1M library Mus musculus clone UUGC2M0141120 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                     AZ842796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0095 row: G column: 1
Contact: Robert B.
University of Utah
                              Unpublished (2000)
Contact: Robert B.
                                                                      plasmid inserts
                                                                                           Mouse whole genome
                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                            Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                  Dunn, D., Aoyagi, A.,
                                                                                                                                                                                                                                                                                                    house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer: CACACAGGAAACAGCTATGACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (g1|473214|gb|RF129072.1), a copy-number inducible derivative of plasmid RL. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          musculus C57BL/6J (male) was obtained
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:10090"
/clone="UUGC1M0095G10"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   clone_lib-"Mouse 10kb plasmid UUGC1M library"
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note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.78;
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       Genome Center
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                                                                                         scaffolding with paired end reads from 10kb
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                                                                                                                                                                                  Barber, M.,
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Pred. No. 1.2e+07
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AI085431
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Best Local Similarity
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                                                                                                                                                                                                          AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               ow82h11.s1 Soares_fetal_liver_spleen_lNFLS_S1 Homo sapiens cDNA
clone IMAGE:1653381 3' similar to TR:Q15409 Q15409 RTVP-1 PROTEIN.
  Onpublished (1975)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
There Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                            AI085431
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Fax: 801 585 7177
                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                             Unpublished (1997)
                                                                                                                                                                             National Cancer Institute,
                                                                                                                                                                                                     NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                       AI085431.1 GI:3423854
                                                                                                                                                       fumor Gene Index
Trace considered overall poor quality
                                                                                                                                                                                                                               Mammalia; Eutheria;
(bases 1 to 28)
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Length: 10000 Std Err
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Strand selected for ampicillin resistance."

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           organism="Mus musculus"
strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sculus C57BL/6J (male) was oboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   one_lib="Mouse 10kb plasmid UUGC1M library"
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one="UUGC2M0141120"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FOW:
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-"Vector: PWD42nv; Purified genomic, DWA from M.
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Pred. No. 1.2e+07
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llin resistance."
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                                                                                                                       and Eco RI sites of the modified pT7T3 vector. went through one round of normalization. Librar
                                                                               constructed by Bento Soares and M.Fatima Bonaldo."
4 c 10 g 6 t
                                                                                                                                                               (Pharmacia), digested with Pac I and cloned into the Pac
                                                                                                                                                                                    double-stranded cDNA was ligated to Eco RI adaptors
                                                                                                                                                                                                    /note-"Organ: Liver and Spleen; V
with a modified polylinker; Site_
This is a subtracted version of t
                                                                                                                                                                                                                                                                                                                                'lab_host="DH10B (ampicil
                                                                                                                                                                                                                                                                                                                                                                        'sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                       organism="Homo sapiens"
db xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                         clone_lib="Soares_fetal_liver_spleen_1NFLS_S1"
                                                                                                                                                                                                                                                                                                                                                                                                                one-"IMAGE:1653381"
    2.78;
                                                                                                                                                                                                                                                                                                                                                 _stage="20 week-post conception fetus"
Score 15.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Std Error: 0.00
                                                                                                                                                                                                                                                         and Spleen; Vector: pT7T3D (Pharmacia)
plinker; Site_1: Pac I; Site_2: Eco RI;
yersion of the original Soares fetal
    DB 9;
                                                                                                                                                                                                                                                                                                                                .lin resistant)  
                                                                                                                                                                                                                                                1st strand cDNA was primed
                  Length
                                                                                                                                            Library
```

밁 Š Matches 76 AGTAACTGTAAGAAAAGCCTGGAGTGC 102 2 AGAMACTTTAGGGCAAGGTTGGAGTGC 28 Conservative 0 Mismatches Indels

Gaps

0

Beat

Local

Similarity

Search completed: May 21, 2003, 07:30:04 Job time: 1444 secs